

Amendments to the Specification:

Please replace the paragraph beginning at page 21, line 24 as with the following amended paragraph:

Functional predictions were made with PFAM (available on the ~~World Wide Web~~ Internet at ~~www~~.pfam.wustl.edu/), which is a Hidden Markov Model based database of families of protein domains. No hits were found in searches using either the *C. elegans* or *M. incognita* polypeptides as queries suggesting that these polypeptides are a novel class of PGM-like enzymes. Protein localization was predicted using the TargetP server (available on the Internet ~~World Wide Web~~ at ~~www~~.cbs.dtu.dk/services/TargetP). The *C. elegans* PGM was predicted to be mitochondrial and the *M. incognita* PGM was predicted to be cytosolic.